

Rebecca Page

List of Publications by Year in descending order

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Version: 2024-02-01

55
papers

3,072
citations

218677

26
h-index

175258

52
g-index

66
all docs

66
docs citations

66
times ranked

5915
citing authors

| # | ARTICLE | IF | CITATIONS |
|----|--|------|-----------|
| 1 | Toxin-antitoxin systems in bacterial growth arrest and persistence. <i>Nature Chemical Biology</i> , 2016, 12, 208-214. | 8.0 | 579 |
| 2 | Targeting the disordered C terminus of PTP1B with an allosteric inhibitor. <i>Nature Chemical Biology</i> , 2014, 10, 558-566. | 8.0 | 294 |
| 3 | Structural basis for protein phosphatase 1 regulation and specificity. <i>FEBS Journal</i> , 2013, 280, 596-611. | 4.7 | 195 |
| 4 | Strategies to maximize heterologous protein expression in <i>Escherichia coli</i> with minimal cost. <i>Protein Expression and Purification</i> , 2007, 51, 1-10. | 1.3 | 190 |
| 5 | Spinophilin directs protein phosphatase 1 specificity by blocking substrate binding sites. <i>Nature Structural and Molecular Biology</i> , 2010, 17, 459-464. | 8.2 | 181 |
| 6 | The Molecular Mechanism of Substrate Engagement and Immunosuppressant Inhibition of Calcineurin. <i>PLoS Biology</i> , 2013, 11, e1001492. | 5.6 | 123 |
| 7 | Expanding the PP2A Interactome by Defining a B56-Specific SLiM. <i>Structure</i> , 2016, 24, 2174-2181. | 3.3 | 117 |
| 8 | Understanding the antagonism of retinoblastoma protein dephosphorylation by PNUTS provides insights into the PP1 regulatory code. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2014, 111, 4097-4102. | 7.1 | 112 |
| 9 | Structural and Functional Analysis of the GADD34:PP1 eIF2 γ Phosphatase. <i>Cell Reports</i> , 2015, 11, 1885-1891. | 6.4 | 107 |
| 10 | Conformational Rigidity and Protein Dynamics at Distinct Timescales Regulate PTP1B Activity and Allostery. <i>Molecular Cell</i> , 2017, 65, 644-658.e5. | 9.7 | 96 |
| 11 | The Molecular Basis for Substrate Specificity of the Nuclear NIPP1:PP1 Holoenzyme. <i>Structure</i> , 2012, 20, 1746-1756. | 3.3 | 70 |
| 12 | The <i>MqsR</i> / <i>MqsA</i> toxin/antitoxin system protects <i>Escherichia coli</i> during bile acid stress. <i>Environmental Microbiology</i> , 2015, 17, 3168-3181. | 3.8 | 55 |
| 13 | Dynamic activation and regulation of the mitogen-activated protein kinase p38. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2018, 115, 4655-4660. | 7.1 | 52 |
| 14 | The Ki-67 and RepoMan mitotic phosphatases assemble via an identical, yet novel mechanism. <i>ELife</i> , 2016, 5, . | 6.0 | 50 |
| 15 | PP1:Tautomycin Complex Reveals a Path toward the Development of PP1-Specific Inhibitors. <i>Journal of the American Chemical Society</i> , 2017, 139, 17703-17706. | 13.7 | 46 |
| 16 | Molecular Investigations of the Structure and Function of the Protein Phosphatase 1 γ Spinophilin γ Inhibitor 2 Heterotrimeric Complex. <i>Biochemistry</i> , 2011, 50, 1238-1246. | 2.5 | 44 |
| 17 | KNL1 Binding to PP1 and Microtubules Is Mutually Exclusive. <i>Structure</i> , 2018, 26, 1327-1336.e4. | 3.3 | 44 |
| 18 | ASPP proteins discriminate between PP1 catalytic subunits through their SH3 domain and the PP1 C-tail. <i>Nature Communications</i> , 2019, 10, 771. | 12.8 | 44 |

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|----|---|------|-----------|
| 19 | NIPP1 maintains EZH2 phosphorylation and promoter occupancy at proliferation-related target genes. <i>Nucleic Acids Research</i> , 2013, 41, 842-854. | 14.5 | 42 |
| 20 | The structures of penicillin-binding protein 4 (PBP4) and PBP5 from Enterococci provide structural insights into β -lactam resistance. <i>Journal of Biological Chemistry</i> , 2018, 293, 18574-18584. | 3.4 | 41 |
| 21 | Regulation of protein phosphatase 1 by intrinsically disordered proteins. <i>Biochemical Society Transactions</i> , 2012, 40, 969-974. | 3.4 | 40 |
| 22 | Investigating the human Calcineurin Interaction Network using the β -LxVP SLiM. <i>Scientific Reports</i> , 2016, 6, 38920. | 3.3 | 39 |
| 23 | A dynamic charge-charge interaction modulates PP2A:B56 substrate recruitment. <i>ELife</i> , 2020, 9, . | 6.0 | 37 |
| 24 | Molecular basis for the binding and selective dephosphorylation of Na ⁺ /H ⁺ exchanger 1 by calcineurin. <i>Nature Communications</i> , 2019, 10, 3489. | 12.8 | 36 |
| 25 | Strategies to make protein serine/threonine (PP1, calcineurin) and tyrosine phosphatases (PTP1B) druggable: Achieving specificity by targeting substrate and regulatory protein interaction sites. <i>Bioorganic and Medicinal Chemistry</i> , 2015, 23, 2781-2785. | 3.0 | 33 |
| 26 | Structural and Regulatory Changes in PBP4 Trigger Decreased β -Lactam Susceptibility in <i>Enterococcus faecalis</i> . <i>MBio</i> , 2018, 9, . | 4.1 | 32 |
| 27 | A Quantitative Chemical Proteomic Strategy for Profiling Phosphoprotein Phosphatases from Yeast to Humans. <i>Molecular and Cellular Proteomics</i> , 2018, 17, 2448-2461. | 3.8 | 29 |
| 28 | Identification of the substrate recruitment mechanism of the muscle glycogen protein phosphatase 1 holoenzyme. <i>Science Advances</i> , 2018, 4, eaau6044. | 10.3 | 28 |
| 29 | SDS22 selectively recognizes and traps metal-deficient inactive PP1. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2019, 116, 20472-20481. | 7.1 | 28 |
| 30 | Molecular Insights into the Fungus-Specific Serine/Threonine Protein Phosphatase Z1 in <i>Candida albicans</i> . <i>MBio</i> , 2016, 7, . | 4.1 | 22 |
| 31 | The KIM-family protein-tyrosine phosphatases use distinct reversible oxidation intermediates: Intramolecular or intermolecular disulfide bond formation. <i>Journal of Biological Chemistry</i> , 2017, 292, 8786-8796. | 3.4 | 21 |
| 32 | Redox Regulation of a Gain-of-Function Mutation (N308D) in SHP2 Noonan Syndrome. <i>ACS Omega</i> , 2017, 2, 8313-8318. | 3.5 | 19 |
| 33 | PP2A/B56 substrate recruitment as defined by the retinoblastoma-related protein p107. <i>ELife</i> , 2021, 10, . | 6.0 | 19 |
| 34 | BdcA, a Protein Important for <i>Escherichia coli</i> Biofilm Dispersal, Is a Short-Chain Dehydrogenase/Reductase that Binds Specifically to NADPH. <i>PLoS ONE</i> , 2014, 9, e105751. | 2.5 | 18 |
| 35 | Leveraging New Definitions of the LxVP SLiM To Discover Novel Calcineurin Regulators and Substrates. <i>ACS Chemical Biology</i> , 2019, 14, 2672-2682. | 3.4 | 17 |
| 36 | Structure-Guided Exploration of SDS22 Interactions with Protein Phosphatase PP1 and the Splicing Factor BCLAF1. <i>Structure</i> , 2019, 27, 507-518.e5. | 3.3 | 16 |

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|----|--|------|-----------|
| 37 | Cooperative dynamics across distinct structural elements regulate PTP1B activity. <i>Journal of Biological Chemistry</i> , 2020, 295, 13829-13837. | 3.4 | 16 |
| 38 | The catalytic activity of TCPTP is auto-regulated by its intrinsically disordered tail and activated by Integrin alpha-1. <i>Nature Communications</i> , 2022, 13, 94. | 12.8 | 16 |
| 39 | The structure of the RCAN1:CN complex explains the inhibition of and substrate recruitment by calcineurin. <i>Science Advances</i> , 2020, 6, . | 10.3 | 15 |
| 40 | The mode of action of the Protein tyrosine phosphatase 1B inhibitor Ertiprotafib. <i>PLoS ONE</i> , 2020, 15, e0240044. | 2.5 | 15 |
| 41 | NMR Spectroscopy to Study MAP Kinase Binding to MAP Kinase Phosphatases. <i>Methods in Molecular Biology</i> , 2016, 1447, 181-196. | 0.9 | 14 |
| 42 | Interaction of Kinase-Interaction-Motif Protein Tyrosine Phosphatases with the Mitogen-Activated Protein Kinase ERK2. <i>PLoS ONE</i> , 2014, 9, e91934. | 2.5 | 13 |
| 43 | NMR Based SARS-CoV-2 Antibody Screening. <i>Journal of the American Chemical Society</i> , 2021, 143, 7930-7934. | 13.7 | 10 |
| 44 | A peculiar IclR family transcription factor regulates para-hydroxybenzoate catabolism in <i>Streptomyces coelicolor</i> . <i>Nucleic Acids Research</i> , 2018, 46, 1501-1512. | 14.5 | 9 |
| 45 | Structures of Dynamic Protein Complexes: Hybrid Techniques to Study MAP Kinase Complexes and the ESCRT System. <i>Methods in Molecular Biology</i> , 2018, 1688, 375-389. | 0.9 | 9 |
| 46 | Strategies for Improving Crystallization Success Rates. <i>Methods in Molecular Biology</i> , 2008, 426, 345-362. | 0.9 | 9 |
| 47 | The interaction of p38 with its upstream kinase MKK6. <i>Protein Science</i> , 2021, 30, 908-913. | 7.6 | 7 |
| 48 | The structure of SDS22 provides insights into the mechanism of heterodimer formation with PP1. <i>Acta Crystallographica Section F, Structural Biology Communications</i> , 2018, 74, 817-824. | 0.8 | 5 |
| 49 | Degradation of the <i>E. coli</i> antitoxin MqsA by the proteolytic complex ClpXP is regulated by zinc occupancy and oxidation. <i>Journal of Biological Chemistry</i> , 2022, 298, 101557. | 3.4 | 5 |
| 50 | ¹ H, ¹⁵ N and ¹³ C sequence specific backbone assignment of the vanadate inhibited hematopoietic tyrosine phosphatase. <i>Biomolecular NMR Assignments</i> , 2018, 12, 5-9. | 0.8 | 4 |
| 51 | Oxidative stress promotes fibrosis in systemic sclerosis through stabilization of a kinase-phosphatase complex. <i>JCI Insight</i> , 2022, 7, . | 5.0 | 3 |
| 52 | Preparation of Phosphorylated Proteins for NMR Spectroscopy. <i>Methods in Enzymology</i> , 2019, 614, 187-205. | 1.0 | 2 |
| 53 | ¹ H, ¹⁵ N and ¹³ C sequence specific backbone assignment of the MAP kinase binding domain of the dual specificity phosphatase 1 and its interaction with the MAPK p38. <i>Biomolecular NMR Assignments</i> , 2021, 15, 243-248. | 0.8 | 0 |
| 54 | Structural biology of MAPK (p38/ERK) regulation by phosphatases and scaffolding proteins. <i>FASEB Journal</i> , 2012, 26, 763.2. | 0.5 | 0 |

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|----|---|-----|-----------|
| 55 | Discovery of Protein Phosphatase 2A Substrates. FASEB Journal, 2018, 32, 795.2. | 0.5 | 0 |